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 ISSN: 0027-8424.
- AU Moore T (Reprint); Constancia M; Zubair M; Bailleul B; Feil R; Sasaki H; Reik W

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- GA The Genuine Article (R) Number: YF393
- TI Multiple imprinted sense and antisense transcripts, differential methylation and tandem repeats in a putative imprinting control region upstream of mouse Igf2
- AU Moore T (Reprint); Constancia M; Zubair M; Bailleul B; Feil R; Sasaki H; Reik W
- CS BABRAHAM INST, DEPT GENET & DEV, CAMBRIDGE CB2 4AT, ENGLAND (Reprint); KYUSHU UNIV, INST GENET INFORMAT, HIGASHI KU, FUKUOKA 812, JAPAN
- CYA ENGLAND; JAPAN
- SO PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF

AMERICA, (11 NOV 1997) Vol. 94, No. 23, pp. 12509-12514.

ISSN: 0027-8424.

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- LIFE FS
- English LA
- REC Reference Count: 28 ED
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- Proceedings of the National Academy of Sciences of the United States of SO America (1997), 94(23), 12509-12514 CODEN: PNASA6; ISSN: 0027-8424
- Moore, T.; Constancia, M.; Zubair, M.; Bailleul, B.; Feil, R.; AU Sasaki, H.; Reik, W.

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     Journal of cell science, (1997 Apr) 110 ( Pt 7) 801-7.
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     Zilberfarb V; Pietri-Rouxel F; Jockers R; Krief S; Delouis C;
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     conformational change of constitutive receptor dimers.
     Journal of Biological Chemistry, (18 Jul 2003) Vol. 278, No. 29, pp.
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     Nucleic acid compositions, kits, and methods for identification,
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Clancy, Brian; Pittman, Debra M.

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WO 2001046697 A2 **20010628** PIPATENT NO. WO 2001046697 ΡI WO 2001046697 HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG Antisense oligonucleotides ITRL: THU (Therapeutic use); BIOL (Biological study); USES (Uses) (nucleic acid compns., kits, and methods for identification, assessment, prevention, and therapy of human breast cancer) 391840-26-9, DNA (human gene DPP-I) 391840-95-2, DNA (human isolate IT391841-60-4, DNA (human gene UCPH cDNA) 391842-52-7 Korean cDNA) 391842-53-8, DNA (human clone 323380 cDNA) 391843-44-0 391844-54-5 391844-55-6, DNA (human gene BTF5 cDNA) 391845-63-9, DNA (human bikunin 391845-65-1, DNA (human gene OPG cDNA) 391847-56-6, DNA (human cDNA) 391849-07-3, DNA (human karyopherin beta 3 391847-74-8 Hlark cDNA) 391850-64-9, DNA (human gene SAP18 cDNA) 391853-22-8, DNA (human cDNA) 391854-60-7, DNA (human gene GPP130 cDNA) 391853-26-2 gene fb19) 391996-11-5, DNA (human tyrosyl-tRNA 391995-75-8 391854-61-8 synthetase cDNA) 391996-17-1, DNA (human gene RETL1 cDNA) 391997-17-4 392000-38-3, DNA (human clone RP3-434P1) 392007-47-5, DNA 391998-11-1 (human gene hCTR1 cDNA) 392009-65-3 392009-82-4, DNA (human clone hsalg7 cDNA) 392011-03-9, DNA (human gene OB-RGRP)

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STIC-Biotech/ChemLib

From:

Wollenberger, Louis V.

Sent:

Thursday, August 04, 2005 1:49 PM

To: Subject:

STIC-Biotech/ChemLib Sequence search request

August 4, 2005

Hi:

Re: Patent Application 10/774721 (Jockers et al.)

Please search the following sequences:

1. A score overlength search of nucleic acid sequence SEQ ID NO:21, looking for oligos 10 to 60 nucleotides in length that are at least 70% identical to a sequence in SEQ ID NO:21.

- 2. A standard search of nucleic acid sequence SEQ ID NO:21 against the nucleic acid databases.
- 3. A length limited search of oligonucleotide sequence SEQ ID NO:37, looking for sequences having at least 60% identity with SEQ ID NO:37.
- 4. A length limited search of oligonucleotide sequence SEQ ID NO:38, looking for sequences having at least 60% identity with SEQ ID NO:38.

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